

Qy	540	IYABDLYIYhrLeuArg1ProArgLaserProGlu1leGluAspSerAlaAsnG	560
Db	2005	GTAATGGGAGCTAGTTCAGTTCCCGGCCAGAGTCCAGGATGCATTG	2064
Qy	560	IYSerArgArgLeuMetLeuProProSerThrProThrProSerGlyProProA	580
Db	2065	GTCRGGCAGCTACGTACCAACCCCTCATACCCACTCCCTGCGCCCTCGGA	2124
Qy	580	RGGIYAlaGluSerValHisSerPheTYHISalaAspCysHisLeuGluProValAlaG	600
Db	2125	GGGGTGGAGGTCTACAGCTCTACAGCTCTACAGCTCTACAGCTCTACAGCT	2184
Qy	600	YSGInAlaProProArgCYSProSerGluAlaSerGlyArgThrValGlySerGlyL	620
Db	2185	GCCAGGCACCCCTCCAGATGCCATCGAGGATCTGAGGATCTGAGGATCGGGGA	2244
Qy	620	YsvAltyrProThrValHisthrSerProProGluLeuLysAspLysAlaLeuV	640
Db	2245	AGGTGACCCACTGTGATACAGCCCTCACAGAGATACTGAGGATAAACGCTAG	2304
Qy	640	AlGluValAlaProSerProGlyProProGlyLeuProProGlyLysSer	660
Db	2305	TGGAGCTGCCCTCACGCTCTCTGAGAACACGAGTACGGGGCTGCTGCATNGCTCT	2364
Qy	660	RophsSerSerMechIlysLeuLeuGluLysSerThrGlyAlaCysHISerSerC	680
Db	2365	CCTTCAGCTCATGGAACTGGCTCAGCTCCACATCCACCTTCACATCCACCTGGGC	2424
Qy	680	YsLybIlesSerProCysSerLysAlaAspSerGlyAlaCysGlyProAspSerCysBP	700
Db	2425	GCAAAATCTCGGCCCTGTCAGGAGAACAGTGAGCTGGGCCAGATGTCAGTGTGIC	2484
Qy	700	RotyrosAlaArgThrGlyAlaGlyGluProGluSerAlaAspHisValMetProAspS	720
Db	2485	CCTACTGTGCCGGACAGGGCAAGGGCAGACTCCGAGATGGATGAGATCTGGTGCACAC	2544
Qy	720	ErabsSerGluAlaValtyrGluPheThrClnAspAlaGlnHisSerAspLeuArgAspP	740
Db	2545	CAGACAGCGGGCTCTGTAGACTGTCACAGGGCGTACAGGAGCTGACTCCGGGATC	2604
Qy	740	RohIsserArgArgGlnArgSerLeuGluProSerValLeuA	760
Db	2605	CCCAGGCCGCGGGCACCGGACACTCTGTCACACTGTCGACATGGCTGAGCTGCTGG	2664
Qy	760	LaPheTrpArgLysCysAspThrPheArgLysSilevalAspSerLysTyrPheGlyA	780
Db	2665	CTTTCGGAGGCTGATCTGACATTCGGAAAGATCGTAACTTGCCGAAATCTGGCC	2724
Qy	780	RGGIYIleMetIleAlaAlleLeuValAsnThrLeuSerMetGlyIleGlutYHISgluG	800
Db	2725	GGGGAAATCATGATCCCCTCATCTGGTAAATCACTCAGCATGGCATCGAGTACACGAGC	2784
Qy	800	InProGluGluLeuThrAspAlaLeuGluLysSerAsnIleValIpheThrSerLeuPheA	820
Db	2785	AGCCCAAGGGCTCACAAAGCCCTGAAATCAGAACATGCTTACAGGCTCTTCG	2844
Qy	820	IaLeuLysMetLysLeuLeuValValValleSerValTrpGluIleValGlyGlnGlnG	860
Db	2845	CCTTGAGATGCTGCAAATGCTTACGGCTTACGGTCACTGTCATGTCATGTCATG	2904
Qy	840	YAsnIlePheAspGlyValleValleValleSerValTrpGluIleValGlyGlnGlnG	880
Db	2905	ACAACATCTCTGATCTGTCATGTCATGTCATGTCATGTCATGTCATGTCATGTCATG	2964
Qy	860	LysGlyLeuSerValLeuArgLysLeuValValleSerValTrpGluIleValGlyGlnGlnG	900
Db	2965	GAGGTGGCTGTTGGCTGGGACTTCGGCTGATGCGGCTGCTGAGCTGGTGGCTGCT	3024
Qy	880	HeLeuProAlaLeuGlnArgGlnIleuValValleLeuMetIshThrMetAspAsnValAlaT	900
Db	3025	TCCCTGGCCCTCCACACCTGGCTCAGGTCTAGGGCTATAGGGTGTGGGCTGAGCTGAGCCTGCA	3084
Qy	900	HrphectysMetLeuLeuMetLeuPhaIlePheSerIleLeuGlyMetHisLeuP	920

